

SEQUENCE LISTING

```
<110> Axxima Pharmaceuticals AG
           Schubart, Daniel
 5
           Habenberger, Peter
           Stein-Gerlach, Matthias
           Bevec, Dorian
    <120> Cellular Kinases Involved in Cytomegalovirus Infection and their
10
    Inhibition
    <130> AXM-004.1 US
    <140> 09/981,397
15
    <141> 2001-10-16
    <150> 60/240,750
    <151> 2000-10-16
20
    <160> 22
    <170> PatentIn version 3.1
    <210> 1
25
    <211> 18
    <212> DNA
    <213> Artificial Sequence
    <220>
30
    <223> synthetic polyT primer
    <220>
    <221> misc_feature
    <222>
           (18)..(18)
35
    <223> n = a,c,g, or t
    <400> 1
    tttttttt tttttvn
                                                                          18
40
    <210> 2
    <211>
           28
    <212> DNA
45
    <213> Artificial Sequence
    <223> cDNA probe for NIK-interacting kinase
50
    <400> 2
    gtcctggagg gctctttttg atgaaacc
                                                                          28
    <210> 3
55
    <211> 30
    <212> DNA
    <213> Artificial Sequence
    <220>
60
    <223> cDNA probe for cellular protein RIP
```

```
<400> 3
    gtgctcaatg cagttgggcc ccttgtacac
                                                                          30
5
    <210> 4
    <211> 27
    <212> DNA
    <213> Artificial Sequence
10
    <220>
    <223> cDNA probe for cellular protein kinase RICK
    <400> 4
15
                                                                          27
    gtcgagcagc ggagtgtgga tgtgcag
    <210> 5
    <211> 23
    <212> PRT
20
    <213> Artificial Sequence
    <220>
    <223> NIK peptide immunogen
25
    <400> 5
    Cys Asn Pro Thr Asn Thr Arg Pro Gln Ser Asp Thr Pro Glu Ile Arg
30
    Lys Tyr Lys Lys Arg Phe Asn
                20
35
    <210> 6
          34
    <211>
    <212> DNA
    <213> Artificial Sequence
40
    <223>
          loxP sequence for recombination vector
    <400> 6
45
    ataacttcgt atagcataca ttatacgaag ttat
                                                                          34
    <210> 7
    <211> 31
50
    <212> DNA
    <213> Artificial Sequence
    <220>
    <223> primer US-10(200900)SpeI
55
    <400> 7
    gctcactagt ggcctagcct ggctcatggc c
                                                                          31
```

11

60

<210> 8

```
<211> 33
    <212> DNA
    <213> Artificial Sequence
 5 <220>
    <223> primer US-10(198918)PacI
    <400> 8
    gtccttaatt aagacgtggt tgtggtcacc gaa
                                                                         33
10
    <210> 9
    <211> 33
    <212> DNA
    <213> Artificial Sequence
    <220>
    <223> primer US-9-3'PmeI
20
   <400> 9
    ctcggtttaa acgacgtgag gcgctccgtc acc
                                                                         33
    <210> 10
25
    <211> 31
    <212> DNA
    <213> Artificial Sequence
    <220>
30
   <223> primer US-5'CLAI
    <400> 10
    ttgcatcgat acggtgtgag ataccacgat g
                                                                         31
35
    <210> 11
    <211> 20
    <212> DNA
    <213> Artificial Sequence
40
    <220>
    <223> primer US-9(198789)
    <400> 11
45
   tgacgcgagt attacgtgtc
                                                                         20
    <210> 12
    <211> 19
50
    <212> DNA
    <213> Artificial Sequence
    <220>
    <223> primer US-10(199100)
55
    <400> 12
    ctcctcctga tatgcggtt
                                                                         19
60
    <210> 13
```

q

<211> 2501

<212> DNA

<213> Homo sapiens

<400> 13 ggcaccagtc tctagaaaag aagtcagctc tggttcggag aagcagcggc tggcgtgggc 60 catccgggga atgggcgccc tcgtgaccta gtgttgcggg gcaaaaaggg tcttgccggc 120 10 ctcgctcgtg caggggcgta tctgggcgcc tgagcgcgca gtgggagcct tgggagccgc 180 cgcagcaggg ggcacacccg gaaccggcct gagcgcccgg gaccatgaac ggggaggcca 240 tetgeagege cetgeceace attecetace acaaactege egacetgege tacetgagee 300 15 geggegeete tggcaetgtg tegteegeee gecaegeaga etggegegte caggtggeeg 360 tgaagcacct gcacatccac actccgctgc tcgacagtga aagaaaggat gtcttaagag 420 20 aagctgaaat tttacacaaa gctagattta gttacattct tccaattttg ggaatttgca 480 atgageetga atttttggga atagttaetg aatacatgee aaatggatea ttaaatgaae 540 tectacatag gaaaactgaa tateetgatg ttgettggee attgagattt egeateetge 600 25 atgaaattgc ccttggtgta aattacctgc acaatatgac tcctccttta cttcatcatg 660 acttgaagac tcagaatatc ttattggaca atgaatttca tgttaagatt gcagattttg 720 30 gtttatcaaa gtggcgcatg atgtccctct cacagtcacg aagtagcaaa tctgcaccag 780 aaggagggac aattatctat atgccacctg aaaactatga acctggacaa aaatcaaggg 840 ccagtatcaa gcacgatata tatagctatg cagttatcac atgggaagtg ttatccagaa 900 35 aacagccttt tgaagatgtc accaatcctt tgcagataat gtatagtgtg tcacaaggac 960 ategaeetgt tattaatgaa gaaagtttge catatgatat aceteaeega geaegtatga 1020 40 tctctctaat agaaagtgga tgggcacaaa atccagatga aagaccatct ttcttaaaat 1080 gtttaataga acttgaacca gttttgagaa catttgaaga gataactttt cttgaagctg 1140 ttattcagct aaagaaaaca aagttacaga gtgtttcaag tgccattcac ctatgtgaca 1200 45 agaagaaaat ggaattatct ctgaacatac ctgtaaatca tggtccacaa gaggaatcat 1260 gtggatcctc tcagctccat gaaaatagtg gttctcctga aacttcaagg tccctgccag 1320 50 ctcctcaaga caatgatttt ttatctagaa aagctcaaga ctgttatttt atgaagctgc 1380 atcactgtcc tggaaatcac agttgggata gcaccatttc tggttctcaa agggctgcat 1440 tetgtgatea caagaceact ceatgetett cageaataat aaateeacte teaáetgeag 1500 55 gaaactcaga acgtctgcag cctggtatag cccagcagtg gatccagagc aaaagggaag 1560 acattgtgaa ccaaatgaca gaagcctgcc ttaaccagtc gctagatgcc cttctgtcca 1620 60 gggacttgat catgaaagag gactatgaac ttgttagtac caagcctaca aggacctcaa 1680

aagtcagaca attactagac actactgaca tccaaggaga agaatttgcc aaagttatag 1740 tacaaaaatt gaaagataac aaacaaatgg gtcttcagcc ttacccggaa atacttgtgg 1800 5 tttctagatc accatcttta aatttacttc aaaataaaag catgtaagtg actgtttttc 1860 aagaagaaat gtgtttcata aaaggatatt tatatctctg ttgctttgac tttttttata 1920 10 taaaatccgt gagtattaaa gctttattga aggttctttg ggtaaatatt agtctccctc 1980 catgacactg cagtattttt tttaattaat acaagtaaaa agttgaattt ggttgaattt 2040 gctacatagt tcaattttta tgtctctttt gttaacagaa accactttta aaggatagta 2100 15 attattcttg tttataacag tgccttaagg tatgatgtat ttctgatgga agccattttc 2160 acattcatgt tcttcatgga ttatttgtta cttgtctaag atgcaatttg attttatgaa 2220 20 gtatataccc tttacccacc agagacagta cagaatccct gccctaaaat cccaggctta 2280 attgccctac aaagggttat taatttaaaa ctccattatt aggattacat tttaaagttt 2340 tatttatgaa ttccctttaa aaatgatatt tcaaaggtaa aacaatacaa tataaagaaa 2400 25 aaaataaata tattaatacc ggcttcctgt ccccattttt aacctcagcc ttccctactg 2460 tcaccaacaa ccaagctaaa taaagtcaac agcctgatgt g 2501 30 14 <210> <211> 1080 <212> PRT Homo sapiens <213> 35 <400> 14 Met Asn Gly Glu Ala Ile Cys Ser Ala Leu Pro Thr Ile Pro Tyr His 40 Lys Leu Ala Asp Leu Arg Tyr Leu Ser Arg Gly Ala Ser Gly Thr Val 20 45 Ser Ser Ala Arg His Ala Asp Trp Arg Val Gln Val Ala Val Lys His 35 4.0 50 Leu His Ile His Thr Pro Leu Leu Asp Ser Glu Arg Lys Asp Val Leu 50 55 Arg Glu Ala Glu Ile Leu His Lys Ala Arg Phe Ser Tyr Ile Leu Pro 55 80 65 Ile Leu Gly Ile Cys Asn Glu Pro Glu Phe Leu Gly Ile Val Thr Glu

60

	Tyr	Met	Pro	Asn 100	Gly	Ser	Leu	Asn	Glu 105	Leu	Leu	His	Arg	Lys 110	Thr	Glu
5	Tyr	Pro	Asp 115	Val	Ala	Trp	Pro	Leu 120	Arg	Phe	Arg	Ile	Leu 125	His	Glu	Ile
10	Ala	Leu 130	Gly	Val	Asn	Tyr	Leu 135	His	Asn	Met	Thr	Pro 140	Pro	Leu	Leu	His
15	His 145	Asp	Leu	Lys	Thr	Gln 150	Asn	Ile	Leu	Leu	Asp 155	Asn	Glu	Phe	His	Val 160
20	Lys	Ile	Ala	Asp	Phe 165	Gly	Leu	Ser	Lys	Trp 170	Arg	Met	Met	Ser	Leu 175	Ser
	Gln	Ser	Arg	Ser 180	Ser	Lys	Ser	Ala	Pro 185	Glu	Gly	Gly	Thr	Ile 190	Ile	Tyr
25	Met	Pro	Pro 195	Glu	Asn	Tyr	Glu	Pro 200	Gly	Gln	Lys	Ser	Arg 205	Ala	Ser	Ile
30	Lys	His 210	Asp	Ile	Tyr	Ser	Tyr 215	Ala	Val	Ile	Thr	Trp 220	Glu	Val	Leu	Ser
35	Arg 225	Lys	Gln	Pro	Phe	Glu 230	Asp	Val	Thr	Asn	Pro 235	Leu	Gln	Ile	Met	Tyr 240
40	Ser	Val	Ser	Gln	Gly 245	His	Arg	Pro	Val	Ile 250	Asn	Glu	Glu	Ser	Leu 255	Pro
	Tyr	Asp	Ile	Pro 260	His	Arg	Ala	Arg	Met 265	Ile	Ser	Leu	Ile	Glu 270	Ser	Gly
45	Trp	Ala	Gln 275	Asn	Pro	Asp	Glu	Arg 280	Pro	Ser	Phe	Leu	Lys 285	Cys	Leu	Ile
50	Glu	Leu 290	Glu	Pro	Val	Leu	Arg 295	Thr	Phe	Glu	Glu	Ile 300	Thr	Phe	Leu	Glu
55	Ala 305	Val	Ile	Gln	Leu	Lys 310	Lys	Thr	Lys	Leu	Gln 315	Ser	Val	Ser	Ser	Ala 320
60	Ile	His	Leu	Cys	Asp 325	Lys	Lys	Lys	Met	Glu 330	Leu	Ser	Leu	Asn	11e 335	Pro

	Val	Asn	His	Gly 340	Pro	Gln	Glu	Glu	Ser 345	Сув	Gly	Ser	Ser	Gln 350	Leu	His
5	Glu	Asn	Ser 355	Gly	Ser	Pro	Glu	Thr 360	Ser	Arg	Ser	Leu	Pro 365	Ala	Pro	Gln
10	Asp	Asn 370	Asp	Phe	Leu	Ser	Arg 375	Lys	Ala	Gln	Asp	Cys 380	Tyr	Phe	Met	Lys
15	Leu 385	His	His	Cys	Pro	Gly 390	Asn	His	Ser	Trp	Asp 395	Ser	Thr	Ile	Ser	Gly 400
20	Ser	Gln	Arg	Ala	Ala 405	Phe	Cys	Asp	His	Lys 410	Thr	Thr	Pro	Cys	Ser 415	Ser
	Ala	Ile	Ile	Asn 420	Pro	Leu	Ser	Thr	Ala 425	Gly	Asn	Ser	Glu	Arg 430	Leu	Gln
25	Pro	Gly	Ile 435	Ala	Gln	Gln	Trp	Ile 440	Gln	Ser	Lys	Arg	Glu 445	Asp	Ile	Val
30	Asn	Gln 450	Met	Thr	Glu	Ala	Cys 455	Leu	Asn	Gln	Ser	Leu 460	Asp	Ala	Leu	Leu
35	Ser 465	Arg	Asp	Leu	Ile	Met 470	Lys	Glu	Asp	Tyr	Glu 475	Leu	Val	Ser	Thr	Lys 480
40	Pro	Thr	Arg	Thr	Ser 485	Lys	Val	Arg	Gln	Leu 490	Leu	Asp	Thr	Thr	Asp 495	Ile
	Gln	Gly	Glu	Glu 500	Phe	Ala	Lys	Val	Ile 505		Gln	Lys	Leu	Lys 510	Asp	Asn
45	Lys	Gln	Met 515	Gly	Leu	Gln	Pro	Tyr 520	Pro	Glu	Ile	Leu	Val 525	Val	Ser	Arg
50	Ser	Pro 530	Ser	Leu	Asn	Leu	Leu 535	Gln	Asn	Lys	Ser	Met 540	Met	Asn	Gly	Glu
55	Ala 545	Ile	Cys	Ser	Ala	Leu 550	Pro	Thr	Ile	Pro	Tyr 555	His	Lys	Leu	Ala	Asp 560
60	Leu	Arg	Tyr	Leu	Ser 565	Arg	Gly	Ala	Ser	Gly 570	Thr	Val	Ser	Ser	Ala 575	Arg

1)

	His	Ala	Asp	Trp 580	Arg	Val	Gln	Val	Ala 585	Val	Lys	His	Leu	His 590	Ile	His
5	Thr	Pro	Leu 595	Leu	Asp	Ser	Glu	Arg 600	Lys	Asp	Val	Leu	Arg 605	Glu	Ala	Glu
10	Ile	Leu 610	His	Lys	Ala	Arg	Phe 615	Ser	Tyr	Ile	Leu	Pro 620	Ile	Leu	Gly	Ile
15	Cys 625	Asn	Glu	Pro	Glu	Phe 630	Leu	Gly	Ile	Val	Thr 635	Glu	Tyr	Met	Pro	Asn 640
20	Gly	Ser	Leu	Asn	Glu 645	Leu	Leu	His	Arg	Lys 650	Thr	Glu	Tyr	Pro	Asp 655	Val
	Ala	Trp	Pro	Leu 660	Arg	Phe	Arg	Ile	Leu 665	His	Glu	Ile	Ala	Leu 670	Gly	Val
25	Asn	Tyr	Leu 675	His	Asn	Met	Thr	Pro 680	Pro	Leu	Leu	His	His 685	Asp	Leu	Lys
30	Thr	Gln 690	Asn	Ile	Leu	Leu	Asp 695	Asn	Glu	Phe	His	Val 700	Lys	Ile	Ala	Asp
35	Phe 705	Gly	Leu	Ser	Lys	Trp 710	Arg	Met	Met	Ser	Leu 715	Ser	Gln	Ser	Arg	Ser 720
40	Ser	Lys	Ser	Ala	Pro 725	Glu	Gly	Gly	Thr	Ile 730	Ile	Tyr	Met	Pro	Pro 735	Glu
	Asn	Tyr	Glu	Pro 740	Gly	Gln	Lys	Ser	Arg 745	Ala	Ser	Ile	Lys	His 750	Asp	Ile
45	Tyr	Ser	Tyr 755	Ala	Val	Ile	Thr	Trp 760	Glu	Val	Leu	Ser	Arg 765	Lys	Gln	Pro
50	Phe	Glu 770	Asp	Val	Thr	Asn	Pro 775	Leu	Gln	Ile	Met	Tyr 780	Ser	Val	Ser	Gln
55	Gly 785	His	Arg	Pro	Val	Ile 790	Asn	Glu	Glu	Ser	Leu 795	Pro	Tyr	Asp	Ile	Prc 800
60	His	Arg	Ala	Arg	Met 805	Ile	Ser	Leu	Ile	Glu 810	Ser	Gly	Trp	Ala	Gln 815	Asn

	Pro	Asp	Glu	Arg 820	Pro	Ser	Phe	Leu	Lys 825	Cys	Leu	Ile	Glu	Leu 830	Glu	Pro
5	Val	Leu	Arg 835	Thr	Phe	Glu	Glu	Ile 840	Thr	Phe	Leu	Glu	Ala 845	Val	Ile	Gln
10	Leu	Lys 850	Lys	Thr	Lys	Leu	Gln 855	Ser	Val	Ser	Ser	Ala 860	Ile	His	Leu	Cys
15	Asp 865	Lys	Lys	Lys	Met	Glu 870	Leu	Ser	Leu	Asn	Ile 875	Pro	Val	Asn	His	Gly 880
20	Pro	Gln	Glu	Glu	Ser 885	Cys	Gly	Ser	Ser	Gln 890	Leu	His	Glu	Asn	Ser 895	Gly
	Ser	Pro	Glu	Thr 900	Ser	Arg	Ser	Leu	Pro 905	Ala	Pro	Gln	Asp	Asn 910	Asp	Phe
25	Leu	Ser	Arg 915	Lys	Ala	Gln	Asp	Cys 920	-	Phe	Met	Lys	Leu 925	His	His	Cys
30	Pro	Gly 930	Asn	His	Ser	Trp	Asp 935	Ser	Thr	Ile	Ser	Gly 940	Ser	Gln	Arg	Ala
35	Ala 945	Phe	Cys	Asp	His	Lys 950	Thr	Thr	Pro	Cys	Ser 955	Ser	Ala	Ile	Ile	Asn 960
40	Pro	Leu	Ser	Thr	Ala 965	Gly	Asn	Ser	Glu	Arg 970	Leu	Gln	Pro	Gly	Ile 975	Ala
	Gln	Gln	Trp	Ile 980	Gln	Ser	Lys	Arg	Glu 985	Asp	Ile	Val	Asn	Gln 990	Met	Thr
45	Glu	Ala	Cys 995	Leu	Asn	Gln	Ser	Leu 1000		o Ala	a Lei	ı Leı	1 Ser		cg As	sp Leu
50	Ile	Met 1010		s Glu	ı As <u>r</u>	р Туз	c Glu 101		∋u Va	al Se	er Th		/s I 020	?ro :	Thr <i>l</i>	Arg
55	Thr	Ser 1025	-	s Val	l Arg	g Glı	101		eu As	sp Th	ır Tl		sp 3	[le (Gln (Gly
60	Glu	Glu 1040		e Ala	a Lys	s Val	l Ile 104		al G	ln Ly	/s Le		/s /	Asp A	Asn l	Cys

```
Gln Met Gly Leu Gln Pro Tyr Pro Glu Ile Leu Val Val Ser Arg
                              1060
        1055
                                                   1065
5
    Ser Pro Ser Leu Asn Leu Leu Gln Asn Lys Ser Met
        1070
                              1075
                                                   1080
10
    <210>
           15
           2617
    <211>
     <212>
           DNA
     <213> Homo sapiens
15
    <220>
     <221> misc_feature
     <222>
           (2141)..(2141)
     <223> n = a,c,g or t
20
    <220>
    <221> misc_feature
    <222>
           (2311)..(2311)
    <223> n = a,c,q or t
25
    <220>
    <221> misc feature
    <222>
           (2452)..(2452)
30
    <223> n = a,c,g or t
    <220>
    <221> misc_feature
35
    <222>
           (2496)..(2496)
    <223> n = a,c,g or t
    <400> 15
40
    atgcaaccag acatgtcctt gaatgtcatt aagatgaaat ccagtgactt cctggagagt
                                                                           60
    gcagaactgg acagcggagg ctttgggaag gtgtctctgt gtttccacag aacccaggga
                                                                          120
    ctcatgatca tgaaaacagt gtacaagggg cccaactgca ttgagcacaa cgaggccctc
                                                                          180
45
    ttggaggagg cgaagatgat gaacagactg agacacagcc gggtggtgaa gctcctgggc
                                                                          240
    gtcatcatag aggaagggaa gtactccctg gtgatggagt acatggagaa gggcaacctg
                                                                          300
50
    atgcacgtgc tgaaagccga gatgagtact ccgctttctg taaaaggaag gataattttg
                                                                          360
    gaaatcattg aaggaatgtg ctacttacat ggaaaaggcg tgatacacaa ggacctgaag
                                                                          420
    cctgaaaata tccttgttga taatgacttc cacattaaga tcgcagacct cggccttgcc
                                                                          480
55
    tcctttaaga tgtggagcaa actgaataat gaagagcaca atgagctgag ggaagtggac
                                                                          540
    ggcaccgcta agaagaatgg cggcaccctc tactacatgg cgcccgagca cctgaatgac
                                                                          600
60
    gtcaacgcaa agcccacaga gaagteggat gtgtacagct ttgctgtagt actctgggcg
                                                                          660
```

atatttgcaa ataaggagcc atatgaaaat gctatctgtg agcagcagtt gataatgtgc 720 ataaaatctg ggaacaggcc agatgtggat gacatcactg agtactgccc aagagaaatt 780 5 atcagtetea tgaagetetg etgggaageg aatceggaag eteggeegae attteetgge 840 900 attgaagaaa aatttaggcc tttttattta agtcaattag aagaaagtgt agaagaggac 10 gtgaagagtt taaagaaaga gtattcaaac gaaaatgcag ttgtgaagag aatgcagtct 960 cttcaacttg attgtgtggc agtaccttca agccggtcaa attcagccac agaacagcct 1020 ggttcactgc acagttccca gggacttggg atgggtcctg tggaggagtc ctggtttgct 1080 15 ccttccctgg agcacccaca agaagagaat gagcccagcc tgcagagtaa actccaagac 1140 gaagccaact accatcttta tggcagccgc atggacaggc agacgaaaca gcagcccaga 1200 20 cagaatgtgg cttacaacag agaggaggaa aggagacgca gggtctccca tgaccctttt 1260 gcacagcaaa gaccttacga gaattttcag aatacagagg gaaaaggcac tgtttattcc 1320 agtgcagcca gtcatggtaa tgcagtgcac cagccctcag ggctcaccag ccaacctcaa 1380 25 gtactgtatc agaacaatgg attatatagc tcacatggct ttggaacaag accactggat 1440 ccaggaacag caggtcccag agtttggtac aggccaattc caagtcatat gcctagtctq 1500 30 cataatatcc cagtgcctga gaccaactat ctaggaaata cacccaccat gccattcagc 1560 teettgeeac caacagatga atetataaaa tataccatat acaatagtae tggcatteag 1620 attggagcct acaattatat ggagattggt gggacgagtt catcactact agacagcaca 1680 35 aatacgaact tcaaagaaga gccagctgct aagtaccaag ctatctttga taataccact 1740 agtctgacgg ataaacacct ggacccaatc agggaaaatc tgggaaagca ctggaaaaac 1800 40 tgtgcccgta aactgggctt cacacagtct cagattgatg aaattgacca tgactatgag 1860 cgagatggac tgaaagaaaa ggtttaccag atgctccaaa agtgggtgat gagggaaggc 1920 ataaagggag ccacggtggg gaagctggcc caggcgctcc accagtgttc caggatcgac 1980 45 cttctgagca gcttgattta cgtcagccag aactaaccct ggatgggcta cggcagctga 2040 agtggacgcc tcacttagcg gataacccca gaaagttggc tgcctcagag cattcagaat 2100 50 tetgteetea etgatagggg ttetgtgtet geagaaattt ngttteetgt aetteatage 2160 tggagaatgg ggaaagaaat ctgcagcaaa ggggtctcac tctgttgcca ggctggtctc 2220 aaacttetgg actcaagtga teeteeegee teggeettee aaagtgetgg gatateagge 2280 55 actgagccac tgcgcccagt caacaatccg ntctgaggaa agcgtaagca ggaagacctc 2340 ttaatggcat agcaccaata aaaaaatgac tcctagttgt gtttggaaag ggagagaaga 2400 60 gatgtctgag gaaggtcatg ttctttcagc ttatggcatt tcctagagtt tngttgaagc 2460

aagaagaaaa actcagagaa tataaaatca actttnaaaa ttgtgtgctc tcttcttcac gtaggctcct gttaaaaaca aagtgcagtc agattctaag ccctgttcag agacttcgcg gatcacagct gcagctcacc gccacatcac aggatcc <210> 16 <211> 1342 <212> PRT <213> Homo sapiens <400> 16 Met Gln Pro Asp Met Ser Leu Asn Val Ile Lys Met Lys Ser Ser Asp Phe Leu Glu Ser Ala Glu Leu Asp Ser Gly Gly Phe Gly Lys Val Ser Leu Cys Phe His Arg Thr Gln Gly Leu Met Ile Met Lys Thr Val Tyr Lys Gly Pro Asn Cys Ile Glu His Asn Glu Ala Leu Leu Glu Glu Ala Lys Met Met Asn Arg Leu Arg His Ser Arg Val Val Lys Leu Leu Gly Val Ile Ile Glu Glu Gly Lys Tyr Ser Leu Val Met Glu Tyr Met Glu Lys Gly Asn Leu Met His Val Leu Lys Ala Glu Met Ser Thr Pro Leu Ser Val Lys Gly Arg Ile Ile Leu Glu Ile Ile Glu Gly Met Cys Tyr Leu His Gly Lys Gly Val Ile His Lys Asp Leu Lys Pro Glu Asn Ile Leu Val Asp Asn Asp Phe His Ile Lys Ile Ala Asp Leu Gly Leu Ala Ser Phe Lys Met Trp Ser Lys Leu Asn Asn Glu Glu His Asn Glu Leu

Arg Glu Val Asp Gly Thr Ala Lys Lys Asn Gly Gly Thr Leu Tyr Tyr

180 185 190

5	Met	Ala	Pro 195	Glu	His	Leu	Asn	Asp 200	Val	Asn	Ala	Lys	Pro 205	Thr	Glu	Lys
10	Ser	Asp 210	Val	Tyr	Ser	Phe	Ala 215	Val	Val	Leu	Trp	Ala 220	Ile	Phe	Ala	Asn
	Lys 225	Glu	Pro	Tyr	Glu	Asn 230	Ala	Ile	Cys	Glu	Gln 235	Gln	Leu	Ile	Met	Cys 240
15	Ile	Lys	Ser	Gly	Asn 245	Arg	Pro	Asp	Val	Asp 250	Asp	Ile	Thr	Glu	Tyr 255	Cys
20	Pro	Arg	Glu	Ile 260	Ile	Ser	Leu	Met	Lys 265	Leu	Cys	Trp	Glu	Ala 270	Asn	Pro
25	Glu	Ala	Arg 275	Pro	Thr	Phe	Pro	Gly 280	Ile	Glu	Glu	Lys	Phe 285	Arg	Pro	Phe
30	Tyr	Leu 290	Ser	Gln	Leu	Glu	Glu 295	Ser	Val	Glu	Glu	Asp 300	Val	Lys	Ser	Leu
	Lys 305	Lys	Glu	Tyr	Ser	Asn 310	Glu	Asn	Ala	Val	Val 315	Lys	Arg	Met	Gln	Ser 320
35	Leu	Gln	Leu	Asp	Cys 325	Val	Ala	Val	Pro	Ser 330	Ser	Arg	Ser	Asn	Ser 335	Ala
40	Thr	Glu	Gln	Pro 340	Gly	Ser	Leu	His	Ser 345	Ser	Gln	Gly	Leu	Gly 350	Met	Gly
45	Pro	Val	Glu 355	Glu	Ser	Trp		Ala 360		Ser	Leu	Glu	His 365	Pro	Gln	Glu
50	Glu	Asn 370	Glu	Pro	Ser	Leu	Gln 375	Ser	Lys	Leu	Gln	Asp 380	Glu	Ala	Asn	Tyr
	His 385	Leu	Tyr	Gly	Ser	Arg 390	Met	Asp	Arg	Gln	Thr 395	Lys	Gln	Gln	Pro	Arg 400
55	Gln	Asn	Val	Ala	Tyr 405	Asn	Arg	Glu	Glu	Glu 410	Arg	Arg	Arg	Arg	Val 415	Ser
60	His	Asp	Pro	Phe	Ala	Gln	Gln	Arg	Pro	Tyr	Glu	Asn	Phe	Gln	Asn	Thr

420 425 430

5	Glu	Gly	Lys 435	Gly	Thr	Val	Tyr	Ser 440	Ser	Ala	Ala	Ser	His 445	Gly	Asn	Ala
10	Val	His 450	Gln	Pro	Ser	Gly	Leu 455	Thr	Ser	Gln	Pro	Gln 460	Val	Leu	Tyr	Gln
	Asn 465	Asn	Gly	Leu	Tyr	Ser 470	Ser	His	Gly	Phe	Gly 475	Thr	Arg	Pro	Leu	Asp 480
15	Pro	Gly	Thr	Ala	Gly 485	Pro	Arg	Val	Trp	Tyr 490	Arg	Pro	Ile	Pro	Ser 495	His
20	Met	Pro	Ser	Leu 500	His	Asn	Ile	Pro	Val 505	Pro	Glu	Thr	Asn	Tyr 510	Leu	Gly
25	Asn	Thr	Pro 515	Thr	Met	Pro	Phe	Ser 520	Ser	Leu	Pro	Pro	Thr 525	Asp	Glu	Ser
30	Ile	Lys 530	Tyr	Thr	Ile	Tyr	Asn 535	Ser	Thr	Gly	Ile	Gln 540	Ile	Gly	Ala	Tyr
	Asn 545	Tyr	Met	Glu	Ile	Gly 550	Gly	Thr	Ser	Ser	Ser 555	Leu	Leu	Asp	Ser	Thr 560
35	Asn	Thr	Asn	Phe	Lys 565	Glu	Glu	Pro	Ala	Ala 570	Lys	Tyr	Gln	Ala	Ile 575	Phe
40	Asp	Asn	Thr	Thr 580	Ser	Leu	Thr	Asp	Lys 585	His	Leu	Asp	Pro	Ile 590	Arg	Glu
45	Asn	Leu	Gly 595	Lys	His	Trp	Lys	Asn 600	Cys	Ala	Arg	Lys	Leu 605	Gly	Phe	Thr
50	Gln	Ser 610	Gln	Ile	Asp	Glu	Ile 615	Asp	His	Asp	Tyr	Glu 620	Arg	Asp	Gly	Leu
	Lys 625	Glu	Lys	Val	Tyr	Gln 630	Met	Leu	Gln	Lys	Trp 635	Val	Met	Arg	Glu	Gly 640
55	Ile	Lys	Gly	Ala	Thr 645	Val	Gly	Lys	Leu	Ala 650	Gln	Ala	Leu	His	Gln 655	Cys
60	Ser	Arg	Ile	Asp	Leu	Leu	Ser	Ser	Leu	Ile	Tyr	Val	Ser	Gln	Asn	Met

660 665 670

5	Gln	Pro	Asp 675	Met	Ser	Leu	Asn	Val 680	Ile	Lys	Met	Lys	Ser 685	Ser	Asp	Phe
10	Leu	Glu 690	Ser	Ala	Glu	Leu	Asp 695	Ser	Gly	Gly	Phe	Gly 700	Lys	Val	Ser	Leu
	Cys 705	Phe	His	Arg	Thr	Gln 710	Gly	Leu	Met	Ile	Met 715	Lys	Thr	Val	Tyr	Lys 720
15	Gly	Pro	Asn	Cys	Ile 725	Glu	His	Asn	Glu	Ala 730	Leu	Leu	Glu	Glu	Ala 735	Lys
20	Met	Met	Asn	Arg 740	Leu	Arg	His	Ser	Arg 745	Val	Val	Lys	Leu	Leu 750	Gly	Val
25	Ile	Ile	Glu 755	Glu	Gly	Lys	Tyr	Ser 760	Leu	Val	Met	Glu	Tyr 765	Met	Glu	Lys
30	Gly	Asn 770	Leu	Met	His	Val	Leu 775	Lys	Ala	Glu	Met	Ser 780	Thr	Pro	Leu	Ser
50	Val 785	Lys	Gly	Arg	Ile	Ile 790	Leu	Glu	Ile	Ile	Glu 795	Gly	Met	Сув	Tyr	Leu 800
35	His	Gly	Lys	Gly	Val 805	Ile	His	Lys	Asp	Leu 810	Lys	Pro	Glu	Asn	Ile 815	Leu
40	Val	Asp	Asn	Asp 820	Phe	His	Ile	Lys	Ile 825	Ala	Asp	Leu	Gly	Leu 830	Ala	Ser
45	Phe	Lys	Met 835	Trp	Ser	Lys	Leu	Asn 840	Asn	Glu	Glu	His	Asn 845	Glu	Leu	Arg
50	Glu	Val 850	Asp	Gly	Thr	Ala	Lys 855	Lys	Asn	Gly	Gly	Thr 860	Leu	Tyr	Tyr	Met
50	Ala 865	Pro	Glu	His	Leu	Asn 870	Asp	Val	Asn	Ala	Lys 875	Pro	Thr	Glu	Lys	Ser 880
55	Asp	Val	Tyr	Ser	Phe 885	Ala	Val	Val	Leu	Trp 890	Ala	Ile	Phe	Ala	Asn 895	Lys
60	Glu	Pro	Tyr	Glu	Asn	Ala	Ile	Cys	Glu	Gln	Gln	Leu	Ile	Met	Cys	Ile

900 905 910

5	Lys	Ser	Gly 915	Asn	Arg	Pro	Asp	Val 920	Asp	Asp	Ile 7	Fhr Gl 92		r Cys	s Pro
10	Arg	Glu 930	Ile	Ile	Ser	Leu	Met 935	Lys	Leu	Cys	-	Glu Al 940	la Ası	n Pro	Glu
	Ala 945	Arg	Pro	Thr	Phe	Pro 950	Gly	Ile	Glu	Glu	Lys I 955	Phe Ar	g Pro	⊃ Phe	960
15	Leu	Ser	Gln	Leu	Glu 965	Glu	Ser	Val	Glu	Glu 970	Asp '	Val Ly	rs Se	c Let 975	-
20	Lys	Glu	Tyr	Ser 980	Asn	Glu	Asn	Ala	Val 985	Val	Lys A	Arg Me	et Gli 990		Leu
25	Gln	Leu	Asp 995	Cys	Val	Ala	Val	Pro 1000		s Sei	r Arg		Asn :	Ser <i>P</i>	Ala Thr
30	Glu	Gln 1010		Gly	Ser	Leu	His		er Se	er G	ln Gly	y Leu 1020	_	Met	Gly
	Pro	Val 1025		ı Glu	ı Ser	Trp	Phe 103		la Pı	co Se	er Let	ı Glu 1035		Pro	Gln
35	Glu	Glu 1040		ı Glu	Pro	Ser	Let 104		ln S€	er Ly	ys Leı	ı Gln 1050	_	Glu	Ala
40	Asn	Tyr 1055		Leu	ı Tyr	Gly	7 Sei 106		rg Me	et A:	sp Arg	g Gln 1065		Lys	Gln
45	Gln	Pro 1070	-	g Glr	n Asr	ı Val	. Ala		yr As	sn Ai	rg Glı	ı Glu 1080		Arg	Arg
50	Arg	Arg 1085		Ser	His	a Asp) Pro		ne Al	la G	ln Glr	n Arg 1095		Tyr	Glu
	Asn	Phe 1100		n Asn	ı Thr	: Glu	110		ys Gl	ly Tì	nr Val	l Tyr 1110		Ser	Ala
55	Ala	Ser 1115		s Gly	Asr	n Ala	112		is G	ln Pi	ro Sei	r Gly 1125		Thr	Ser
60	Gln	Pro	Glr	ı Val	. Leu	і Туг	Glı	n As	sn As	sn G	ly Lei	ı Tyr	Ser	Ser	His

1130	1135	1140
1130	1133	1140

5	Gly	Phe 1145	Gly	Thr	Arg	Pro	Leu 1150		Pro	Gly	Thr	Ala 1155	Gly	Pro	Arg
10	Val	Trp 1160	Tyr	Arg	Pro	Ile	Pro 1165	Ser	His	Met	Pro	Ser 1170	Leu	His	Asn
	Ile	Pro 1175	Val	Pro	Glu	Thr	Asn 1180	Tyr	Leu	Gly	Asn	Thr 1185	Pro	Thr	Met
15	Pro	Phe 1190	Ser	Ser	Leu	Pro	Pro 1195	Thr	Asp	Glu	Ser	Ile 1200	Lys	Tyr	Thr
20	Ile	Tyr 1205	Asn	Ser	Thr	Gly	Ile 1210	Gln	Ile	Gly	Ala	Tyr 1215	Asn	Tyr	Met
25	Glu	Ile 1220	Gly	Gly	Thr	Ser	Ser 1225	Ser	Leu	Leu	Asp	Ser 1230	Thr	Asn	Thr
30	Asn	Phe 1235	Lys	Glu	Glu	Pro	Ala 1240	Ala	Lys	Tyr	Gln	Ala 1245	Ile	Phe	Asp
	Asn	Thr 1250	Thr	Ser	Leu	Thr	Asp 1255	Lys	His	Leu	Asp	Pro 1260	Ile	Arg	Glu
35	Asn	Leu 1265	Gly	Lys	His	Trp	Lys 1270	Asn	Cys	Ala	Arg	Lys 1275	Leu	Gly	Phe
40	Thr	Gln 1280	Ser	Gln	Ile	Asp	Glu 1285	Ile	Asp	His	Asp	Tyr 1290	Glu	Arg	Asp
45	_	Leu 1295	_	Glu	Lys		Tyr 1300		Met	Leu		Lys 1305	_	Val	Met
50	Arg	Glu 1310	Gly	Ile	Lys	Gly	Ala 1315	Thr	Val	Gly	Lys	Leu 1320	Ala	Gln	Ala
	Leu	His 1325	Gln	Cys	Ser	Arg	Ile 1330	Asp	Leu	Leu	Ser	Ser 1335	Leu	Ile	Tyr
55	Val	Ser 1340	Gln	Asn											

<210> 17

<211> 4596 <212> DNA

<213> Homo sapiens

5 <400> 17 aagcggggga ctgtgccgtg tggaacgtgt agctgttgag aggtggactc tgttaccatt 60 gaggatgttt ggaggatgag tatgtgtggc agaggcacac ataaacaggc agagaccctt 120 10 tgcccctgcc tttctccccc aacccaaggc tgacctgtgt tctcccaggt ctgggattct 180 aagtgacctg ctctgtgttt ggtctctctc aggatgagca caagcctggg agatggcagt 240 gatggaaatg gcctgcccag gtgcccctgg ctcagcagtg gggcagcaga aggaactccc 300 15 caagccaaag gagaagacgc cgccactggg gaagaaacag agctccgtct acaagcttga 360 ggccgtggag aagagccctg tgttctgcgg aaagtgggag atcctgaatg acgtgattac 420 20 caagggcaca gccaaggaag gctccgaggc agggccagct gccatctcta tcatcgccca 480 ggctgagtgt gagaatagcc aagagttcag ccccaccttt tcagaacgca ttttcatcgc 540 tgggtccaaa cagtacagcc agtccgagag tcttgatcag atccccaaca atgtggccca 600 25 tgctacagag ggcaaaatgg cccgtgtgtg ttggaaggga aagcgtcgca gcaaagcccg 660 gaagaaacgg aagaagaaga gctcaaagtc cctggctcat gcaggagtgg ccttggccaa 720 30 accectecce aggacectg ageaggagag etgeaceate eeagtgeagg aggatgagte 780 tccactcggc gccccatatg ttagaaacac cccgcagttc accaagcctc tgaaggaacc 840 aggccttggg caactctgtt ttaagcagct tggcgagggc ctacggccgg ctctgcctcg 900 35 atcagaactc cacaaactga tcagcccctt gcaatgtctg aaccacgtgt ggaaactgca 960 ccaccccag gacggaggcc ccctgccct gcccacgcac cccttcccct atagcagact 1020 40 gcctcatccc ttcccattcc accetctcca gccctggaaa cctcaccctc tggagtcctt 1080 cctgggcaaa ctggcctgtg tagacagcca gaaacccttg cctgacccac acctgagcaa 1140 actggcctgt gtagacagtc caaagcccct gcctggccca cacctggagc ccagctgcct 1200 45 gtctcgtggt gcccatgaga agttttctgt ggaggaatac ctagtgcatg ctctgcaagg 1260 cagegtgage teaagecagg cecacageet gaccageetg gecaagaeet gggeageaeg 1320 50 gggctccaga tcccgggagc ccagccccaa aactgaggac aacgagggtg tcctgctcac 1380 tgagaaactc aagccagtgg attatgagta ccgagaagaa gtccactggg ccacgcacca 1440 gctccgcctg ggcagaggct ccttcggaga ggtgcacagg atggaggaca agcagactgg 1500 55 cttccagtgc gctgtcaaaa aggtgcggct ggaagtattt cgggcagagg agctgatggc 1560 atgtgcagga ttgacctcac ccagaattgt ccctttgtat ggagctgtga gagaagggcc 1620 60 1680 ttgggtcaac atcttcatgg agctgctgga aggtggctcc ctgggccagc tggtcaagga

geagggetgt eteccagagg accgggeeet gtactacetg ggeeaggeee tggagggtet 1740 ggaatacete caeteaegaa ggattetgea tggggaegte aaagetgaea aegtgeteet 1800 5 gtccagcgat gggagccacg cagccctctg tgactttggc catgctgtgt gtcttcaacc 1860 tgatggcctg ggaaagtcct tgctcacagg ggactacatc cctggcacag agacccacat 1920 10 ggctccggag gtggtgctgg gcaggagctg cgacgccaag gtggatgtct ggagcagctg 1980 ctgtatgatg ctgcacatgc tcaacggctg ccacccctgg actcagttct tccgagggcc 2040 getetgeete aagattgeea gegageetee geetgtgagg gagateeeae eeteetgege 2100 15 ccctctcaca gcccaggcca tccaagaggg gctgaggaaa gagcccatcc accgcgtgtc 2160 tgcagcggag ctgggaggga aggtgaaccg ggcactacag caagtgggag gtctgaagag 2220 20 cccttggagg ggagaatata aagaaccaag acatccaccg ccaaatcaag ccaattacca 2280 ccagaccete catgeceage egagagaget ttegecaagg geeceaggge eeeggeeage 2340 tgaggagaca acaggcagag cccctaagct ccagceteet eteccaccag ageceecaga 2400 25 gccaaacaag tctcctccct tgactttgag caaggaggag tctgggatgt gggaaccctt 2460 acctctgtcc tccctggagc cagcccctgc cagaaacccc agctcaccag agcggaaagc 2520 30 aaccgtcccg gagcaggaac tgcagcagct ggaaatagaa ttattcctca acagcctgtc 2580 ccagccattt tctctggagg agcaggagca aattctctcg tgcctcagca tcgacagcct 2640 ctccctgtcg gatgacagtg agaagaaccc atcaaaggcc tctcaaagct cgcgggacac 2700 35 cctgagctca ggcgtacact cctggagcag ccaggccgag gctcgaagct ccagctggaa 2760 catggtgctg gcccgggggc ggcccaccga caccccaagc tatttcaatg gtgtgaaagt 2820 40 ccaaatacag tctcttaatg gtgaacacct gcacatccgg gagttccacc gggtcaaagt 2880 gggagacatc gccactggca tcagcagcca gatcccagct gcagccttca gcttggtcac 2940 caaagacggg cagcctgttc gctacgacat ggaggtgcca gactcgggca tcgacctgca 3000 45 3060 gtgcacactg gcccctgatg gcagcttcgc ctggagctgg agggtcaagc atggccagct ggagaacagg coctaaccot gccotocaco gccggotoca cactgoogga aagcagoott 3120 50 cctgctcggt gcacgatgct gccctgaaaa cacaggctca gccgttccca ggggattgcc 3180 agccccccgg ctcacagtgg gaaccagggc ctcgcagcag caaggtgggg gcaagcagaa 3240 tgcctcccag gatttcacac ctgagccctg cccaccctg ctgaaaaaac atccgccacg 3300 55 tgaagagaca gaaggaggat ggcaggagtt acctggggaa acaaaacagg gatctttttc 3360 tgcccctgct ccagtcgagt tggcctgacc cgcttggatc agtgaccatt tgttggcaga 3420 60 caggggagag cagcttccag cctgggtcag aaggggtggg cgagcccttc ggcccctcac 3480

	cctccaggct	gctgtgagag	tgtcaagtgt	gtaagggccc	aaactcaggt	tcagtgcaga	3540
5	accaggtcag	caggtatgcc	cgcccgtagg	ttaagggggc	cctctaaacc	ccttgcctgg	3600
J	cctcacctgg	ccagctcacc	ccttttgggt	gtaggggaaa	agaatgcctg	accctgggaa	3660
	ggctccctgg	tagaatacac	cacacttttc	aggttgttgc	aacacaggtc	ctgagttgac	3720
10	ctctggttca	gccaaggacc	aaagaaggtg	tgtaagtgaa	gtggttctca	gtccccagac	3780
	atgtgcccct	ttgctgctgg	ctaccactct	tccccagagc	agcaggcccc	gagccccttc	3840
15	aggcccagca	ctgccccaga	ctcgctggca	ctcagttccc	tcatctgtaa	aggtgaaggg	3900
13	tgatgcagga	tatgcctgac	aggaacagtc	tgtggatgga	catgatcagt	gctaaggaaa	3960
	gcagcagaga	gagacgtccg	gcgccccagc	cccactatca	gtgtccagcg	tgctggttcc	4020
20	ccagagcaca	gctcagcatc	acactgacac	tcaccctgcc	ctgcccctgg	ccagagggta	4080
	ctgccgacgg	cactttgcac	tctgatgacc	tcaaagcact	ttcatggctg	ccctctggca	4140
25	gggcagggca	gggcagtgac	actgtaggag	catagcaagc	caggagatgg	ggtgaaggga	4200
23	cacagtcttg	agctgtccac	atgcatgtga	ctcctcaaac	ctcttccaga	tttctctaag	4260
	aatagcaccc	ccttccccat	tgccccagct	tagcctcttc	tcccagggga	gctactcagg	4320
30	actcacgtag	cattaaatca	gctgtgaatc	gtcagggggt	gtctgctagc	ctcaacctcc	4380
	tggggcaggg	gacgccgaga	ctccgtggga	gaagctcatt	cccacatctt	gccaagacag	4440
35	cctttgtcca	gctgtccaca	ttgagtcaga	ctgctcccgg	ggagagagcc	ccggccccca	4500
55	gcacataaag	aactgcagcc	ttggtactgc	agagtctggg	ttgtagagaa	ctctttgtaa	4560
	gcaataaagt	ttggggtgat	gacaaatgtt	aaaaaa			4596
40	<210> 18						
	<211> 1894 <212> PRT	1					
45		o sapiens					
15	<400> 18						
	Met Ala Val	l Met Glu Me 5	et Ala Cys	Pro Gly Ala 10	Pro Gly Sen	r Ala Val 15	
50	1	3		10		13	
	Gly Gln Gl	n Lys Glu Le 20	_	Pro Lys Glu 25	Lys Thr Pro	Pro Leu	
55		20	•		30		
33	Gly Lys Lys	s Gln Ser Se	er Val Tyr 1 40	Lys Leu Glu	Ala Val Glu 45	ı Lys Ser	
60	Pro Val Phe	e Cys Gly Ly	ys Trp Glu	Ile Leu Asn	Asp Val Ile	e Thr Lys	

50 55 60

5	Gly 65	Thr	Ala	Lys	Glu	Gly 70.	Ser	Glu	Ala	Gly	Pro 75	Ala	Ala	Ile	Ser	Ile 80
10	Ile	Ala	Gln	Ala	Glu 85	Cys	Glu	Asn	Ser	Gln 90	Glu	Phe	Ser	Pro	Thr 95	Phe
-	Ser	Glu	Arg	Ile 100	Phe	Ile	Ala	Gly	Ser 105	Lys	Gln	Tyr	Ser	Gln 110	Ser	Glu
15	Ser	Leu	Asp 115	Gln	Ile	Pro	Asn	Asn 120	Val	Ala	His	Ala	Thr 125	Glu	Gly	Lys
20	Met	Ala 130	Arg	Val	Cys	Trp	Lys 135	Gly	Lys	Arg	Arg	Ser 140	Lys	Ala	Arg	Lys
25	Lys 145	Arg	Lys	Lys	Lys	Ser 150	Ser	Lys	Ser	Leu	Ala 155	His	Ala	Gly	Val	Ala 160
30	Leu	Ala	Lys	Pro	Leu 165	Pro	Arg	Thr	Pro	Glu 170	Gln	Glu	Ser	Cys	Thr 175	Ile
	Pro	Val	Gln	Glu 180	Asp	Glu	Ser	Pro	Leu 185	Gly	Ala	Pro	Tyr	Val 190	Arg	Asn
35	Thr	Pro	Gln 195	Phe	Thr	Lys	Pro	Leu 200	Lys	Glu	Pro	Gly	Leu 205	Gly	Gln	Leu
40	Cys	Phe 210	Lys	Gln	Leu	Gly	Glu 215	Gly	Leu	Arg	Pro	Ala 220	Leu	Pro	Arg	Ser
45	Glu 225	Leu	His	Lys	Leu	Ile 230	Ser	Pro	Leu	Gln	Cys 235	Leu	Asn	His	Val	Trp 240
50	Lys	Leu	His	His	Pro 245	Gln	Asp	Gly	Gly	Pro 250	Leu	Pro	Leu	Pro	Thr 255	His
	Pro	Phe	Pro	Tyr 260	Ser	Arg	Leu	Pro	His 265	Pro	Phe	Pro	Phe	His 270	Pro	Leu
55	Gln	Pro	Trp 275	Lys	Pro	His	Pro	Leu 280	Glu	Ser	Phe	Leu	Gly 285	Lys	Leu	Ala
60	Cys	Val	Asp	Ser	Gln	Lys	Pro	Leu	Pro	Asp	Pro	His	Leu	Ser	Lys	Leu

290 295 300

5	Ala 305	Cys	Val	Asp	Ser	Pro 310	Lys	Pro	Leu	Pro	Gly 315	Pro	His	Leu	Glu	Pro 320
10	Ser	Cys	Leu	Ser	Arg 325	Gly	Ala	His	Glu	Lys 330	Phe	Ser	Val	Glu	Glu 335	Tyr
,	Leu	Val	His	Ala 340	Leu	Gln	Gly	Ser	Val 345	Ser	Ser	Ser	Gln	Ala 350	His	Ser
15	Leu	Thr	Ser 355	Leu	Ala	Lys	Thr	Trp 360	Ala	Ala	Arg	Gly	Ser 365	Arg	Ser	Arg
20	Glu	Pro 370	Ser	Pro	Lys	Thr	Glu 375	Asp	Asn	Glu	Gly	Val 380	Leu	Leu	Thr	Glu
25	Lys 385	Leu	Lys	Pro	Val	Asp 390	Tyr	Glu	Tyr	Arg	Glu 395	Glu	Val	His	Trp	Ala 400
30	Thr	His	Gln	Leu	Arg 405	Leu	Gly	Arg	Gly	Ser 410	Phe	Gly	Glu	Val	His 415	Arg
	Met	Glu	Asp	Lys 420	Gln	Thr	Gly	Phe	Gln 425	Cys	Ala	Val	Lys	Lys 430	Val	Arg
35	Leu	Glu	Val 435	Phe	Arg	Ala	Glu	Glu 440	Leu	Met	Ala	Cys	Ala 445	Gly	Leu	Thr
40	Ser	Pro 450	Arg	Ile	Val	Pro	Leu 455	Tyr	Gly	Ala	Val	Arg 460	Glu	Gly	Pro	Trp
45	Val 465	Asn	Ile	Phe	Met	Glu 470	Leu	Leu	Glu	Gly	Gly 475	Ser	Leu	Gly	Gln	Leu 480
50	Val	Lys	Glu	Gln	Gly 485	Cys	Leu	Pro	Glu	Asp 490	Arg	Ala	Leu	Tyr	Tyr 495	Leu
	Gly	Gln	Ala	Leu 500	Glu	Gly	Leu	Glu	Tyr 505	Leu	His	Ser	Arg	Arg 510	Ile	Leu
55	His	Gly	Asp 515	Val	Lys	Ala	Asp	Asn 520	Val	Leu	Leu	Ser	Ser 525	Asp	Gly	Ser
60	His	Ala	Ala	Leu	Cys	Asp	Phe	Gly	His	Ala	Val	Cys	Leu	Gln	Pro	Asp

530 535 540

5	Gly 545	Leu	Gly	Lys	Ser	Leu 550	Leu	Thr	Gly	Asp	Tyr 555	Ile	Pro	Gly	Thr	Glu 560
10	Thr	His	Met	Ala	Pro 565	Glu	Val	Val	Leu	Gly 570	Arg	Ser	Cys	Asp	Ala 575	Lys
	Val	Asp	Val	Trp 580	Ser	Ser	Сув	Cys	Met 585	Met	Leu	His	Met	Leu 590	Asn	Gly
15	Cys	His	Pro 595	Trp	Thr	Gln	Phe	Phe 600	Arg	Gly	Pro	Leu	Cys 605	Leu	Lys	Ile
20	Ala	Ser 610	Glu	Pro	Pro	Pro	Val 615	Arg	Glu	Ile	Pro	Pro 620	Ser	Cys	Ala	Pro
25	Leu 625	Thr	Ala	Gln	Ala	Ile 630	Gln	Glu	Gly	Leu	Arg 635	Lys	Glu	Pro	Ile	His 640
30	Arg	Val	Ser	Ala	Ala 645	Glu	Leu	Gly	Gly	Lys 650	Val	Asn	Arg	Ala	Leu 655	Gln
	Gln	Val	Gly	Gly 660	Leu	Lys	Ser	Pro	Trp 665	Arg	Gly	Glu	Tyr	Lys 670	Glu	Pro
35	Arg	His	Pro 675	Pro	Pro	Asn	Gln	Ala 680	Asn	Tyr	His	Gln	Thr 685	Leu	His	Ala
40	Gln	Pro 690	Arg	Glu	Leu	Ser	Pro 695	Arg	Ala	Pro	Gly	Pro 700	Arg	Pro	Ala	Glu
45	Glu 705	Thr	Thr	Gly	Arg	Ala 710	Pro	Lys	Leu	Gln	Pro 715	Pro	Leu	Pro	Pro	Glu 720
50	Pro	Pro	Glu	Pro	Asn 725	Lys	Ser	Pro	Pro	Leu 730	Thr	Leu	Ser	Lys	Glu 735	Glu
	Ser	Gly	Met	Trp 740	Glu	Pro	Leu	Pro	Leu 745	Ser	Ser	Leu	Glu	Pro 750	Ala	Pro
55	Ala	Arg	Asn 755	Pro	Ser	Ser	Pro	Glu 760	Arg	Lys	Ala	Thr	Val 765	Pro	Glu	Gln
60	Glu	Leu	Gln	Gln	Len	Glu	Tle	Glu	Len	Phe	ī.eu	Asn	Ser	Len	Ser	Gln

770 775 780

5	Pro 785	Phe	Ser	Leu	Glu	Glu 790	Gln	Glu	Gln	Ile	Leu 795	Ser	Cys	Leu	Ser	Ile 800
10	Asp	Ser	Leu	Ser	Leu 805	Ser	Asp	Asp	Ser	Glu 810	Lys	Asn	Pro	Ser	Lys 815	Ala
	Ser	Gln	Ser	Ser 820	Arg	Asp	Thr	Leu	Ser 825	Ser	Gly	Val	His	Ser 830	Trp	Ser
15	Ser	Gln	Ala 835	Glu	Ala	Arg	Ser	Ser 840	Ser	Trp	Asn	Met	Val 845	Leu	Ala	Arg
20	Gly	Arg 850	Pro	Thr	Asp	Thr	Pro 855	Ser	Tyr	Phe	Asn	Gly 860	Val	Lys	Val	Gln
25	Ile 865	Gln	Ser	Leu	Asn	Gly 870	Glu	His	Leu	His	Ile 875	Arg	Glu	Phe	His	Arg 880
30	Val	Lys	Val	Gly	Asp 885	Ile	Ala	Thr	Gly	Ile 890	Ser	Ser	Gln	Ile	Pro 895	Ala
	Ala	Ala	Phe	Ser 900	Leu	Val	Thr	Lys	Asp 905	Gly	Gln	Pro	Val	Arg 910	Tyr	Asp
35	Met	Glu	Val 915	Pro	Asp	Ser	Gly	Ile 920	Asp	Leu	Gln	Cys	Thr 925	Leu	Ala	Pro
10	Asp	Gly 930	Ser	Phe	Ala	Trp	Ser 935	Trp	Arg	Val	Lys	His 940	Gly	Gln	Leu	Glu
1 5	Asn 945	Arg	Pro	Met	Ala	Val 950		Glu	Met	Ala	Cys 955		Gly	Ala	Pro	Gly 960
50	Ser	Ala	Val	Gly	Gln 965	Gln	Lys	Glu	Leu	Pro 970	Lys	Pro	Lys	Glu	Lys 975	Thr
	Pro	Pro	Leu	Gly 980	Lys	Lys	Gln	Ser	Ser 985	Val	Tyr	Lys	Leu	Glu 990	Ala	Val
55	Glu	Lys	Ser 995	Pro	Val	Phe	Cys	Gly 1000		: Trp	Glı	ı Ile	Leu 100		sn As	sp Val

 $60\,$ $\,$ Ile Thr $\,$ Lys Gly Thr Ala Lys Glu Gly Ser Glu Ala Gly Pro Ala

		1010					1015					1020			
5	Ala	Ile 1025	Ser	Ile	Ile	Ala	Gln 1030	Ala	Glu	Cys	Glu	Asn 1035	Ser	Gln	Glu
10	Phe	Ser 1040	Pro	Thr	Phe	Ser	Glu 1045	Arg	Ile	Phe	Ile	Ala 1050	Gly	Ser	Lys

- Gln Tyr Ser Gln Ser Glu Ser Leu Asp Gln Ile Pro Asn Asn Val 1055 1060 1065
- 15
 Ala His Ala Thr Glu Gly Lys Met Ala Arg Val Cys Trp Lys Gly
 1070 1075 1080
- 20 Lys Arg Arg Ser Lys Ala Arg Lys Lys Arg Lys Lys Ser Ser 1085 1090 1095
- Lys Ser Leu Ala His Ala Gly Val Ala Leu Ala Lys Pro Leu Pro 25 1100 1105 1110
- Arg Thr Pro Glu Gln Glu Ser Cys Thr Ile Pro Val Gln Glu Asp 1115 1120 1125
 - Glu Ser Pro Leu Gly Ala Pro Tyr Val Arg Asn Thr Pro Gln Phe 1130 1135 1140
- Thr Lys Pro Leu Lys Glu Pro Gly Leu Gly Gln Leu Cys Phe Lys
 1145 1150 1155
- 40 Gln Leu Gly Glu Gly Leu Arg Pro Ala Leu Pro Arg Ser Glu Leu 1160 1165 1170
- His Lys Leu Ile Ser Pro Leu Gln Cys Leu Asn His Val Trp Lys 45 1175 1180 1185
- Leu His His Pro Gln Asp Gly Gly Pro Leu Pro Leu Pro Thr His 1190 1195 1200
 - Pro Phe Pro Tyr Ser Arg Leu Pro His Pro Phe Pro Phe His Pro 1205 1210 1215
- Leu Gln Pro Trp Lys Pro His Pro Leu Glu Ser Phe Leu Gly Lys
 1220 1225 1230
- 60 Leu Ala Cys Val Asp Ser Gln Lys Pro Leu Pro Asp Pro His Leu

1235	1240	1245

5	Ser	Lys 1250	Leu	Ala	Cys	Val	Asp 1255		Pro	Lys	Pro	Leu 1260	Pro	Gly	Pro
10	His	Leu 1265	Glu	Pro	Ser	Cys	Leu 1270	Ser	Arg	Gly	Ala	His 1275	Glu	Lys	Phe
	Ser	Val 1280	Glu	Glu	Tyr	Leu	Val 1285	His	Ala	Leu	Gln	Gly 1290	Ser	Val	Ser
15	Ser	Ser 1295	Gln	Ala	His	Ser	Leu 1300	Thr	Ser	Leu	Ala	Lys 1305	Thr	Trp	Ala
20	Ala	Arg 1310	Gly	Ser	Arg	Ser	Arg 1315	Glu	Pro	Ser	Pro	Lys 1320	Thr	Glu	Asp
25	Asn	Glu 1325	Gly	Val	Leu	Leu	Thr 1330	Glu	Lys	Leu	Lys	Pro 1335	Val	Asp	Tyr
30	Glu	Tyr 1340	Arg	Glu	Glu	Val	His 1345	Trp	Ala	Thr	His	Gln 1350	Leu	Arg	Leu
	Gly	Arg 1355	Gly	Ser	Phe	Gly	Glu 1360	Val	His	Arg	Met	Glu 1365	Asp	Lys	Gln
35	Thr	Gly 1370	Phe	Gln	Cys	Ala	Val 1375	Lys	Lys	Val	Arg	Leu 1380	Glu	Val	Phe
40	Arg	Ala 1385	Glu	Glu	Leu	Met	Ala 1390	Cys	Ala	Gly	Leu	Thr 1395	Ser	Pro	Arg
45	Ile	Val 1400	Pro	Leu	Tyr	Gly	Ala 1405		Arg	Glu	Gly	Pro 1410	Trp	Val	Asn
50	Ile	Phe 1415	Met	Glu	Leu	Leu	Glu 1420	Gly	Gly	Ser	Leu	Gly 1425	Gln	Leu	Val
	Lys	Glu 1430	Gln	Gly	Cys	Leu	Pro 1435	Glu	Asp	Arg	Ala	Leu 1440	Tyr	Tyr	Leu
55	Gly	Gln 1445	Ala	Leu	Glu	Gly	Leu 1450	Glu	Tyr	Leu	His	Ser 1455	Arg	Arg	Ile
60	Leu	His	Gly	Asp	Val	Lys	Ala	Asp	Asn	Val	Leu	Leu	Ser	Ser	Asp

1460	1465	1470
1460	1465	1470

5	Gly	Ser 1475	His	Ala	Ala	Leu	Cys 1480	_	Phe	Gly	His	Ala 1485	Val	Cys	Leu
10	Gln	Pro 1490	Asp	Gly	Leu	Gly	Lys 1495	Ser	Leu	Leu	Thr	Gly 1500	Asp	Tyr	Ile
	Pro	Gly 1505	Thr	Glu	Thr	His	Met 1510	Ala	Pro	Glu	Val	Val 1515	Leu	Gly	Arg
15	Ser	Cys 1520	_	Ala	Lys	Val	Asp 1525		Trp	Ser	Ser	Cys 1530	Cys	Met	Met
20	Leu	His 1535	Met	Leu	Asn	Gly	Cys 1540	His	Pro	Trp	Thr	Gln 1545	Phe	Phe	Arg
25	Gly	Pro 1550	Leu	Суѕ	Leu	Lys	Ile 1555	Ala	Ser	Glu	Pro	Pro 1560	Pro	Val	Arg
30	Glu	Ile 1565	Pro	Pro	Ser	Cys	Ala 1570	Pro	Leu	Thr	Ala	Gln 1575	Ala	Ile	Gln
	Glu	Gly 1580	Leu	Arg	Lys	Glu	Pro 1585	Ile	His	Arg	Val	Ser 1590	Ala	Ala	Glu
35	Leu	Gly 1595	Gly	Lys	Val	Asn	Arg 1600	Ala	Leu	Gln	Gln	Val 1605	Gly	Gly	Leu
40	Lys	Ser 1610	Pro	Trp	Arg	Gly	Glu 1615	Tyr	Lys	Glu	Pro	Arg 1620	His	Pro	Pro
45		Asn 1625				-	His 1630							Pro	Arg
50	Glu	Leu 1640	Ser	Pro	Arg	Ala	Pro 1645	Gly	Pro	Arg	Pro	Ala 1650	Glu	Glu	Thr
	Thr	Gly 1655	Arg	Ala	Pro	Lys	Leu 1660	Gln	Pro	Pro	Leu	Pro 1665	Pro	Glu	Pro
55	Pro	Glu 1670	Pro	Asn	Lys	Ser	Pro 1675	Pro	Leu	Thr	Leu	Ser 1680	Lys	Glu	Glu
60	Ser	Gly	Met	Trp	Glu	Pro	Leu	Pro	Leu	Ser	Ser	Leu	Glu	Pro	Ala

1685 1690 1695

5	Pro	Ala 1700	Arg	Asn	Pro	Ser	Ser 1705	Pro	Glu	Arg	Lys	Ala 1710	Thr	Val	Pro
10	Glu	Gln 1715	Glu	Leu	Gln	Gln	Leu 1720	Glu	Ile	Glu	Leu	Phe 1725	Leu	Asn	Ser
	Leu	Ser 1730	Gln	Pro	Phe	Ser	Leu 1735	Glu	Glu	Gln		Gln 1740	Ile	Leu	Ser
15	Cys	Leu 1745	Ser	Ile	Asp	Ser	Leu 1750		Leu	Ser	Asp	Asp 1755	Ser	Glu	Lys
20	Asn	Pro 1760	Ser	Lys	Ala	Ser	Gln 1765	Ser	Ser	Arg	Asp	Thr 1770	Leu	Ser	Ser
25	Gly	Val 1775	His	Ser	Trp	Ser	Ser 1780	Gln	Ala	Glu	Ala	Arg 1785	Ser	Ser	Ser
30	Trp	Asn 1790	Met	Val	Leu	Ala	Arg 1795	Gly	Arg	Pro	Thr	Asp 1800	Thr	Pro	Ser
	Tyr	Phe 1805	Asn	Gly	Val	Lys	Val 1810	Gln	Ile	Gln	Ser	Leu 1815	Asn	Gly	Glu
35	His	Leu 1820	His	Ile	Arg	Glu	Phe 1825	His	Arg	Val	Lys	Val 1830	Gly	Asp	Ile
40	Ala	Thr 1835	Gly	Ile	Ser	Ser	Gln 1840	Ile	Pro	Ala	Ala	Ala 1845	Phe	Ser	Leu
45		Thr 1850	_	_	_		Pro 1855		_		_	Met 1860		Val	Pro
50	Asp	Ser 1865	Gly	Ile	Asp	Leu	Gln 1870	Cys	Thr	Leu	Ala	Pro 1875	Asp	Gly	Ser
	Phe	Ala 1880	Trp	Ser	Trp	Arg	Val 1885	Lys	His	Gly	Gln	Leu 1890	Glu	Asn	Arg
55															

60 <210> 19

Pro

<211> 2030

<212> DNA

<213> Homo sapiens

5 <400> 19 tggctggcaa tggccttgct gacctcgagc cgggcccacg tggggacctt tggagcacag 60 cctacgatcc tggtgcaagg ccggtggatg cagaggccag tccatatacc acccaggcct 120 10 gcgaggagcg tggtccccac ccatccagcc catatgtgca agtgcccttg acagagagc 180 tggtcatatc catggtgacc atttatgggc cacaacaggt ccccatctgc gcagtgaacc 240 ctgtgctgag caccttgcag acgtgatctt gcttcgtcct gcagcactgt gcggggcagg 300 15 360 aaaatccaag aggaagaagg atctacggat atcctgcatg tccaagccac ccgcacccaa ccccacaccc ccccggaacc tggactcccg gaccttcatc accattggag acagaaactt 420 20 tgaggtggag gctgatgact tggtgaccat ctcagaactg ggccgtggag cctatggggt 480 ggtagagaag gtgcggcacg cccagagcgg caccatcatg gccgtgaagc ggatccgggc 540 caccgtgaac tcacaggagc agaagcggct gctcatggac ctggacatca acatgcgcac 600 25 ggtcgactgt ttctacactg tcaccttcta cggggcacta ttcagagagg gagacgtgtg 660 gatctgcatg gagctcatgg acacatcctt ggacaagttc taccggaagg tgctggataa 720 30 aaacatgaca attccagagg acatccttgg ggagattgct gtgtctatcg tgcgggccct 780 ggagcatctg cacagcaagc tgtcggtgat ccacagagat gtgaagccct ccaatgtcct 840 tatcaacaag gagggccatg tgaagatgtg tgactttggc atcagtggct acttggtgga 900 35 ctctgtggcc aagacgatgg atgccggctg caagccctac atggcccctg agaggatcaa 960 cccagagctg aaccagaagg gctacaatgt caagtccgac gtctggagcc tgggcatcac 1020 40 catgattgag atggccatcc tgcggttccc ttacgagtcc tgggggaccc cgttccagca 1080 gctgaagcag gtggtggagg agccgtcccc ccagctccca gccgaccgtt tctcccccga 1140 gtttgtggac ttcactgctc agtgcctgag gaagaacccc gcagagcgta tgagctacct 1200 45 ggagctgatg gagcacccct tcttcacctt gcacaaaacc aagaagacgg acattgctgc 1260 cttcgtgaag aagatcctgg gagaagactc ataggggctg ggcctcggac cccactccgg 1320 50 ccctccagag ccccacagcc ccatctgcgg gggcagtgct cacccacacc ataagctact 1380 gecatectgg cccagggeat etgggaggaa ecgaggggge tgeteccace tggetetgtg 1440 gcgagccatt tgtcccaagt gccaaagaag cagaccattg gggctcccag ccaggccctt 1500 55 gtcggcccca ccagtgcctc tccctgctgc tcctaggacc cgtctccagc tgctgagatc 1560 ctggactgag ggggcctgga tgccccctgt ggatgctgct gcccctgcac agcaggctgc 1620 60 1680 cagtgcctgg gtggatgggc caccgccttg cccagcctgg atgccatcca agttgtatat

ttttttaatc tctcgactga atggactttg cacactttgg cccagggtgg ccacacctct 1740 atcccggctt tggtgcgggg tacacaagag gggatgagtt gtgtgaatac cccaagactc 1800 5 ccatgaggga gatgccatga gccgcccaag gccttcccct ggcactggca aacagggcct 1860 ctgcggagca cactggctca cccagtcctg cccgccaccg ttatcggtgt cattcacctt 1920 10 tegtgttttt tttaatttat cetetgttga ttttttettt tgetttatgg gtttggettg 1980 tttttcttgc atggtttgga gctgatcgct tctcccccac cccctagggg 2030 15 <210> 20 <211> 636 <212> PRT <213> Homo sapiens 20 <400> 20 Met Ser Lys Pro Pro Ala Pro Asn Pro Thr Pro Pro Arg Asn Leu Asp 25 Ser Arg Thr Phe Ile Thr Ile Gly Asp Arg Asn Phe Glu Val Glu Ala 20 25 30 Asp Asp Leu Val Thr Ile Ser Glu Leu Gly Arg Gly Ala Tyr Gly Val 35 40 Val Glu Lys Val Arg His Ala Gln Ser Gly Thr Ile Met Ala Val Lys 35 50 Arg Ile Arg Ala Thr Val Asn Ser Gln Glu Gln Lys Arg Leu Leu Met 40 Asp Leu Asp Ile Asn Met Arg Thr Val Asp Cys Phe Tyr Thr Val Thr 45 Phe Tyr Gly Ala Leu Phe Arg Glu Gly Asp Val Trp Ile Cys Met Glu 100 50 Leu Met Asp Thr Ser Leu Asp Lys Phe Tyr Arg Lys Val Leu Asp Lys 115 Asn Met Thr Ile Pro Glu Asp Ile Leu Gly Glu Ile Ala Val Ser Ile 55 130 135 Val Arg Ala Leu Glu His Leu His Ser Lys Leu Ser Val Ile His Arg 160 150 145

• *1 .

60

	Asp	Val	Lys	Pro	Ser 165	Asn	Val	Leu	Ile	Asn 170	Lys	Glu	Gly	His	Val 175	Lys
5	Met	Cys	Asp	Phe 180	Gly	Ile	Ser	Gly	Tyr 185	Leu	Val	Asp	Ser	Val 190	Ala	Lys
10	Thr	Met	Asp 195	Ala	Gly	Cys	Lys	Pro 200	Tyr	Met	Ala	Pro	Glu 205	Arg	Ile	Asn
15	Pro	Glu 210	Leu	Asn	Gln	Lys	Gly 215	Tyr	Asn	Val	Lys	Ser 220	Asp	Val	Trp	Ser
20	Leu 225	Gly	Ile	Thr	Met	Ile 230	Glu	Met	Ala	Ile	Leu 235	Arg	Phe	Pro	Tyr	Glu 240
	Ser	Trp	Gly	Thr	Pro 245	Phe	Gln	Gln	Leu	Lys 250	Gln	Val	Val	Glu	Glu 255	Pro
25	Ser	Pro	Gln	Leu 260	Pro	Ala	Asp	Arg	Phe 265	Ser	Pro	Glu	Phe	Val 270	Asp	Phe
30	Thr	Ala	Gln 275	Cys	Leu	Arg	Lys	Asn 280	Pro	Ala	Glu	Arg	Met 285	Ser	Tyr	Leu
35	Glu	Leu 290	Met	Glu	His	Pro	Phe 295	Phe	Thr	Leu	His	Lys 300	Thr	Lys	Lys	Thr
40	Asp 305	Ile	Ala	Ala	Phe	Val 310	Lys	Lys	Ile	Leu	Gly 315	Glu	Asp	Ser	Met	Ser 320
	Lys	Pro	Pro	Ala	Pro 325	Asn	Pro	Thr	Pro	Pro 330		Asn	Leu	Asp	Ser 335	_
45	Thr	Phe	Ile	Thr 340	Ile	Gly	Asp	Arg	Asn 345	Phe	Glu	Val	Glu	Ala 350	Asp	Asp
50	Leu	Val	Thr 355	Ile	Ser	Glu	Leu	Gly 360	Arg	Gly	Ala	Tyr	Gly 365	Val	Val	Glu
55	Lys	Val 370	Arg	His	Ala	Gln	Ser 375	Gly	Thr	Ile	Met	Ala 380	Val	Lys	Arg	Ile
60	Arg 385	Ala	Thr	Val	Asn	Ser 390	Gln	Glu	Gln	Lys	Arg 395	Leu	Leu	Met	Asp	Leu 400

Asp Ile Asn Met Arg Thr Val Asp Cys Phe Tyr Thr Val Thr Phe Tyr 405 410 5 Gly Ala Leu Phe Arg Glu Gly Asp Val Trp Ile Cys Met Glu Leu Met 425 420 10 Asp Thr Ser Leu Asp Lys Phe Tyr Arg Lys Val Leu Asp Lys Asn Met 435 440 Thr Ile Pro Glu Asp Ile Leu Gly Glu Ile Ala Val Ser Ile Val Arq 15 450 455 Ala Leu Glu His Leu His Ser Lys Leu Ser Val Ile His Arg Asp Val 20 Lys Pro Ser Asn Val Leu Ile Asn Lys Glu Gly His Val Lys Met Cys 25 Asp Phe Gly Ile Ser Gly Tyr Leu Val Asp Ser Val Ala Lys Thr Met 30 Asp Ala Gly Cys Lys Pro Tyr Met Ala Pro Glu Arg Ile Asn Pro Glu Leu Asn Gln Lys Gly Tyr Asn Val Lys Ser Asp Val Trp Ser Leu Gly 35 530 Ile Thr Met Ile Glu Met Ala Ile Leu Arg Phe Pro Tyr Glu Ser Trp 545 550 555 560 40 Gly Thr Pro Phe Gln Gln Leu Lys Gln Val Val Glu Glu Pro Ser Pro 45 Gln Leu Pro Ala Asp Arg Phe Ser Pro Glu Phe Val Asp Phe Thr Ala 580 50 Gln Cys Leu Arg Lys Asn Pro Ala Glu Arg Met Ser Tyr Leu Glu Leu 595 600 605 Met Glu His Pro Phe Phe Thr Leu His Lys Thr Lys Lys Thr Asp Ile 55 610 615 620 Ala Ala Phe Val Lys Lys Ile Leu Gly Glu Asp Ser 630 60

<210> 21 <211> 3745 <212> DNA 5 <213> Homo sapiens

<400> 21 gaatteggea egaggeeatt gaateeeagt eetaacagaa gtaetgegaa tettgtggee 60 10 120 tcattctgaa caaaagggat tagagaagaa aaatctcttg atataaggct tgaaagcaag ggcaggcaat cttggttgtg aatattttct gatttttcca gaaatcaagc agaagattga 180 gctgctgatg tcagttaact ctgagaagtc gtcctcttca gaaaggccgg agcctcaaca 240 15 gaaageteet ttagtteete eteeteeace gecaceacea ecaceacege cacetttgee 300 360 agaccccaca cccccggagc cagaggagga gatcctggga tcagatgatg aggagcaaga 20 ggaccctgcg gactactgca aaggtggata tcatccagtg aaaattggag acctcttcaa 420 tggccggtat catgttatta gaaagcttgg atgggggcac ttctctactg tctggctgtg 480 ctgggatatg caggggaaaa gatttgttgc aatgaaagtt gtaaaaagtg cccagcatta 540 25 tacggagaca gccttggatg aaataaaatt gctcaaatgt gttcgagaaa gtgatcccag 600 tgacccaaac aaagacatgg tggtccagct cattgacgac ttcaagattt caggcatgaa 660 30 tgggatacat gtctgcatgg tcttcgaagt acttggccac catctcctca agtggatcat 720 caaatccaac tatcaaggcc tcccagtacg ttgtgtgaag agtatcattc gacaggtcct 780 tcaagggtta gattacttac acagtaagtg caagatcatt catactgaca taaagccgga 840 35 aaatatettg atgtgtgtgg atgatgcata tgtgagaaga atggcagetg ageetgagtg 900 gcagaaagca ggtgctcctc ctccttcagg gtctgcagtg agtacggctc cacagcagaa 960 40 acctatagga aaaatatcta aaaacaaaaa gaaaaaactg aaaaagaaac agaagaggca 1020 ggctgagtta ttggagaagc gcctgcagga gatagaagaa ttggagcgag aagctgaaag 1080 gaaaataata gaagaaaaca tcacctcagc tgcaccttcc aatgaccagg atggcgaata 1140 45 ctgcccagag gtgaaactaa aaacaacagg attagaggag gcggctgagg cagagactgc 1200 aaaggacaat ggtgaagctg aggaccagga agagaaagaa gatgctgaga aagaaaacat 1260 50 tgaaaaagat gaagatgatg tagatcagga acttgcgaac atagacccta cgtggataga 1320 1380 atcacctaaa accaatggcc atattgagaa tggcccattc tcactggagc agcaactgga cgatgaagat gatgatgaag aagactgccc aaatcctgag gaatataatc ttgatgagcc 1440 55 aaatgcagaa agtgattaca catatagcag ctcctatgaa caattcaatg gtgaattgcc 1500 aaatggacga cataaaattc ccgagtcaca gttcccagag ttttccacct cgttgttctc 1560 60 tggatcctta gaacctgtgg cctgcggctc tgtgctttct gagggatcac cacttactga 1620

gcaagaggag agcagtccat cccatgacag aagcagaacg gtttcagcct ccagtactgg 1680 ggatttgcca aaagcaaaaa cccgggcagc tgacttgttg gtgaatcccc tggatccgcg 1740 5 gaatcgagat aaaattagag taaaaattgc tgacctggga aatgcttgtt gggtgcataa 1800 acacttcacg gaagacatcc agacgcgtca gtaccgctcc atagaggttt taataggagc 1860 10 ggggtacagc acccctgcgg acatctggag cacggcgtgt atggcatttg agctggcaac 1920 1980 gggagattat ttgtttgaac cacattctgg ggaagactat tccagagacg aagaccacat agcccacate atagagetge taggeagtat tecaaggeae tttgetetat etggaaaata 2040 15 ttctcgggaa ttcttcaatc gcagaggaga actgcgacac atcaccaagc tgaagccctg 2100 gagcctcttt gatgtacttg tggaaaagta tggctggccc catgaagatg ctgcacagtt 2160 20 tacagatttc ctgatcccga tgttagaaat ggttccagaa aaacgagcct cagctggcga 2220 atgtcggcat ccttggttga attcttagca aattctacca atattgcatt ctgagctagc 2280 aaatgttccc agtacattgg acctaaacgg tgactctcat tctttaacag gattacaagt 2340 25 gagetggett catceteaga cetttatttt getttgaggt aetgttgttt gacattttge 2400 tttttgtgca ctgtgatcct ggggaagggt agtcttttgt cttcagctaa gtagtttact 2460 30 gaccattttc ttctggaaac aataacatgt ctctaagcat tgtttcttgt gttgtgtgac 2520 attcaaatgt catttttttg aatgaaaaat actttcccct ttgtgttttg gcaggttttg 2580 taactattta tgaagaaata ttttagctga gtactatata atttacaatc ttaagaaatt 2640 35 atcaagttgg aaccaagaaa tagcaaggaa atgtacaatt ttatcttctg gcaaagggac 2700 atcattcctg tattatagtg tatgtaaatg caccctgtaa atgttacttt ccattaaata 2760 40 tgggagggg actcaaattt cagaaaagct accaagtctt gagtgctttg tagcctatgt 2820 tgcatgtagc ggactttaac tgctccaagg agttgtgcaa acttttcatt ccataacagt 2880 cttttcacat tggattttaa acaaagtggc tctgggttat aagatgtcat tctctatatg 2940 45 gcactttaaa ggaagaaaag atatgtttct cattctaaaa tatgcattat aatttagcag 3000 tcccatttgt gattttgcat atttttaaaa gtacttttaa agaagagcaa tttcccttta 3060 50 aaaatgtgat ggctcagtac catgtcatgt tgcctcctct gggcgctgta agttaagctc 3120 3180 tacatagatt aaattggaga aacgtgttaa ttgtgtggaa tgaaaaaata catatttt tggaaaagca tgatcatgct tgtctagaac acaaggtatg gtatatacaa tttgcagtgc 3240 55 agtgggcaga atacttctca cagctcaaag ataacagtga tcacattcat tccataggta 3300 getttaegtg tggetacaac aaattttaet agetttttea ttgtetttee atgaaaegaa 3360 60 gttgagaaaa tgattttccc tttgcaggtt gcacacagtt ttgtttatgc atttccttaa 3420

aattaattgt agactccagg atacaaacca tagtaggcaa tacaatttag aatgtaatat atagaggtat attagcctct ttagaagtca gtggattgaa tgtcttttta ttttaaattt tacattcatt aaggtgcctc gtttttgact ttgtccatta acatttatcc atatgccttt gcaataacta gattgtgaaa agctaacaag tgttgtaaca ataatccatt gtttgaggtg cttgcagttg tcttaaaaat taaagtgttt tggttttttt ttttccagaa aaaaaaaaa aaaaaaaaa aaaaaaaatt cctgc <210> 22 <211> <212> PRT <213> Homo sapiens <400> 22 Met Ser Val Asn Ser Glu Lys Ser Ser Ser Glu Arg Pro Glu Pro Gln Gln Lys Ala Pro Leu Val Pro Leu Pro Asp Pro Thr Pro Pro Glu Pro Glu Glu Glu Ile Leu Gly Ser Asp Asp Glu Glu Glu Glu Asp Pro Ala Asp Tyr Cys Lys Gly Gly Tyr His Pro Val Lys Ile Gly Asp Leu Phe Asn Gly Arg Tyr His Val Ile Arg Lys Leu Gly Trp Gly His Phe Ser Thr Val Trp Leu Cys Trp Asp Met Gln Gly Lys Arg Phe Val Ala Met Lys Val Val Lys Ser Ala Gln His Tyr Thr Glu Thr Ala Leu Asp Glu Ile Lys Leu Leu Lys Cys Val Arg Glu Ser Asp Pro Ser Asp Pro Asn Lys Asp Met Val Val Gln Leu Ile Asp Asp Phe Lys Ile Ser Gly Met Asn Gly Ile

His Val Cys Met Val Phe Glu Val Leu Gly His His Leu Leu Lys Trp Ile Ile Lys Ser Asn Tyr Gln Gly Leu Pro Val Arg Cys Val Lys Ser Ile Ile Arg Gln Val Leu Gln Gly Leu Asp Tyr Leu His Ser Lys Cys Lys Ile Ile His Thr Asp Ile Lys Pro Glu Asn Ile Leu Met Cys Val Asp Asp Ala Tyr Val Arg Met Ala Ala Glu Pro Glu Trp Gln Lys Ala Gly Ala Pro Pro Pro Ser Gly Ser Ala Val Ser Thr Ala Pro Gln Gln Lys Pro Ile Gly Lys Ile Ser Lys Asn Lys Lys Lys Leu Lys Lys Lys Gln Lys Arg Gln Ala Glu Leu Leu Glu Lys Arg Leu Gln Glu Ile Glu Glu Leu Glu Arg Glu Ala Glu Arg Lys Ile Ile Glu Glu Asn Ile Thr Ser Ala Ala Pro Ser Asn Asp Gln Asp Gly Glu Tyr Cys Pro Glu Val Lys Leu Lys Thr Thr Gly Leu Glu Glu Ala Ala Glu Ala Glu Thr Ala Lys Asp Asn Gly Glu Ala Glu Asp Gln Glu Glu Lys Glu Asp Ala Glu Lys Glu Asn Ile Glu Lys Asp Glu Asp Asp Val Asp Gln Glu Leu Ala Asn Ile Asp Pro Thr Trp Ile Glu Ser Pro Lys Thr Asn Gly His Ile Glu Asn Gly Pro Phe Ser Leu Glu Gln Gln Leu Asp Asp Glu

Asp Asp Asp Glu Glu Asp Cys Pro Asn Pro Glu Glu Tyr Asn Leu Asp 405 410 5 Glu Pro Asn Ala Glu Ser Asp Tyr Thr Tyr Ser Ser Ser Tyr Glu Gln 420 425 10 Phe Asn Gly Glu Leu Pro Asn Gly Arg His Lys Ile Pro Glu Ser Gln 435 440 Phe Pro Glu Phe Ser Thr Ser Leu Phe Ser Gly Ser Leu Glu Pro Val 15 450 455 460 Ala Cys Gly Ser Val Leu Ser Glu Gly Ser Pro Leu Thr Glu Gln Glu 470 20 Glu Ser Ser Pro Ser His Asp Arg Ser Arg Thr Val Ser Ala Ser Ser 490 25 Thr Gly Asp Leu Pro Lys Ala Lys Thr Arq Ala Ala Asp Leu Leu Val 30 Asn Pro Leu Asp Pro Arg Asn Arg Asp Lys Ile Arg Val Lys Ile Ala Asp Leu Gly Asn Ala Cys Trp Val His Lys His Phe Thr Glu Asp Ile 35 530 Gln Thr Arg Gln Tyr Arg Ser Ile Glu Val Leu Ile Gly Ala Gly Tyr 550 40 Ser Thr Pro Ala Asp Ile Trp Ser Thr Ala Cys Met Ala Phe Glu Leu 565 570 575 45 Ala Thr Gly Asp Tyr Leu Phe Glu Pro His Ser Gly Glu Asp Tyr Ser 580 585 50 Arg Asp Glu Asp His Ile Ala His Ile Ile Glu Leu Leu Gly Ser Ile 595 600 Pro Arg His Phe Ala Leu Ser Gly Lys Tyr Ser Arg Glu Phe Phe Asn 55 610 615 620 Arg Arg Gly Glu Leu Arg His Ile Thr Lys Leu Lys Pro Trp Ser Leu 625 630 635 640 60

Phe Asp Val Leu Val Glu Lys Tyr Gly Trp Pro His Glu Asp Ala Ala Gln Phe Thr Asp Phe Leu Ile Pro Met Leu Glu Met Val Pro Glu Lys Arg Ala Ser Ala Gly Glu Cys Arg His Pro Trp Leu Asn Ser Met Ser Val Asn Ser Glu Lys Ser Ser Ser Ser Glu Arg Pro Glu Pro Gln Gln Pro Pro Leu Pro Asp Pro Thr Pro Pro Glu Pro Glu Glu Glu Ile Leu Gly Ser Asp Asp Glu Glu Glu Glu Asp Pro Ala Asp Tyr Cys Lys Gly Gly Tyr His Pro Val Lys Ile Gly Asp Leu Phe Asn Gly Arg Tyr His Val Ile Arg Lys Leu Gly Trp Gly His Phe Ser Thr Val Trp Leu Cys Trp Asp Met Gln Gly Lys Arg Phe Val Ala Met Lys Val Val Lys Ser Ala Gln His Tyr Thr Glu Thr Ala Leu Asp Glu Ile Lys Leu Leu Lys Cys Val Arg Glu Ser Asp Pro Ser Asp Pro Asn Lys Asp Met Val Val Gln Leu Ile Asp Asp Phe Lys Ile Ser Gly Met Asn Gly Ile His Val Cys Met Val Phe Glu Val Leu Gly His His Leu Leu Lys Trp Ile Ile 850 855 - The state of the stat Lys Ser Asn Tyr Gln Gly Leu Pro Val Arg Cys Val Lys Ser Ile Ile

Arg Gln Val Leu Gln Gly Leu Asp Tyr Leu His Ser Lys Cys Lys Ile Ile His Thr Asp Ile Lys Pro Glu Asn Ile Leu Met Cys Val Asp Asp Ala Tyr Val Arg Arg Met Ala Ala Glu Pro Glu Trp Gln Lys Ala Gly Ala Pro Pro Pro Ser Gly Ser Ala Val Ser Thr Ala Pro Gln Gln Lys Pro Ile Gly Lys Ile Ser Lys Asn Lys Lys Lys Leu Lys Lys Lys Gln Lys Arg Gln Ala Glu Leu Leu Glu Lys Arg Leu Gln Glu Ile Glu Glu Leu Glu Arg Glu Ala Glu Arg Lys Ile Ile Glu Glu Asn Ile Thr Ser Ala Ala Pro Ser Asn Asp Gln Asp Gly Glu Tyr Cys Pro Glu Val Lys Leu Lys Thr Thr Gly Leu Glu Glu Ala Glu Ala Glu Thr Ala Lys Asp Asn Gly Glu Ala Glu Asp Gln Glu Glu Lys Glu Asp Ala Glu Lys Glu Asn Ile Glu Lys Asp Glu Asp Asp Val Asp Gln Glu Leu Ala Asn Ile Asp Pro Thr Trp Ile Glu Ser Pro Lys Thr Asn Gly His Ile Glu Asn Gly Pro Phe Ser Leu Glu Gln Gln Leu Asp Asp Glu Asp Asp Glu Glu Asp Cys Pro Asn Pro Glu Glu 1085 1090 1095 Tyr Asn Leu Asp Glu Pro Asn Ala Glu Ser Asp Tyr Thr Tyr Ser 1100 1105 1110

	Ser	Ser 1115	Tyr	Glu	Gln	Phe	Asn 1120	Gly	Glu	Leu	Pro	Asn 1125	Gly	Arg	His
5	Lys	Ile 1130	Pro	Glu	Ser	Gln	Phe 1135	Pro	Glu	Phe	Ser	Thr 1140	Ser	Leu	Phe
10	Ser	Gly 1145	Ser	Leu	Glu	Pro	Val 1150	Ala	Сув	Gly	Ser	Val 1155	Leu	Ser	Glu
15	Gly	Ser 1160	Pro	Leu	Thr	Glu	Gln 1165	Glu	Glu	Ser	Ser	Pro 1170	Ser	His	Asp
20	Arg	Ser 1175	Arg	Thr	Val	Ser	Ala 1180	Ser	Ser	Thr	Gly	Asp 1185	Leu	Pro	Lys
	Ala	Lys 1190	Thr	Arg	Ala	Ala	Asp 1195	Leu	Leu	Val	Asn	Pro 1200	Leu	Asp	Pro
25	Arg	Asn 1205	Arg	Asp	Lys	Ile	Arg 1210	Val	Lys	Ile	Ala	Asp 1215	Leu	Gly	Asn
30	Ala	Cys 1220	Trp	Val	His	Lys	His 1225	Phe	Thr	Glu	Asp	Ile 1230	Gln	Thr	Arg
35	Gln	Tyr 1235	Arg	Ser	Ile	Glu	Val 1240	Leu	Ile	Gly	Ala	Gly 1245	Tyr	Ser	Thr
40	Pro	Ala 1250	Asp	Ile	Trp	Ser	Thr 1255	Ala	Cys	Met	Ala	Phe 1260	Glu	Leu	Ala
	Thr	Gly 1265	Asp	Tyr	Leu	Phe	Glu 1270	Pro	His	Ser	Gly	Glu 1275	Asp	Tyr	Ser
45	Arg	Asp 1280	Glu	Asp	His	Ile	Ala 1285	His	Ile	Ile	Glu	Leu 1290	Leu	Gly	Ser
50	Ile	Pro 1295	Arg	His	Phe	Ala	Leu 1300	Ser	Gly	Lys	Tyr	Ser 1305	Arg	Glu	Phe
55	Phe	Asn 1310	Arg	Arg	Gly	Glu	Leu 1315	Arg	His	Ile	Thr	Lys 1320	Leu	Lys	Pro
60	Trp	Ser 1325	Leu	Phe	Asp	Val	Leu 1330	Val	Glu	Lys	Tyr	Gly 1335	Trp	Pro	His

Glu Asp Ala Ala Gln Phe Thr Asp Phe Leu Ile Pro Met Leu Glu 1340 1345 1350

5 Met Val Pro Glu Lys Arg Ala Ser Ala Gly Glu Cys Arg His Pro 1355 1360 1365

10 Trp Leu Asn Ser 1370